

PTO/SB/08B (10-01)
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Substitute for form 1449B/PTO		Complete if Known	
		Application Number	09/858.289
INFORMATION DISCLOSURE STATEMENT BY APPLICANT (use as many sheets as necessary)		Filing Date	May 15, 2001
		First Named Inventor	Nicholas Schork, et al.
		Group Art Unit	Unknown 1631
		Examiner Name	Unassigned
		Attorney Docket Number	55.US4.DIV
Sheet	2	of	3

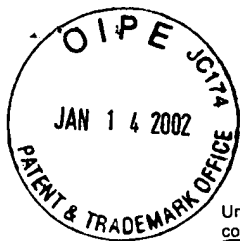
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Examiner Initials	Cite No. 1	Include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item (book, magazine, journal, serial, symposium, catalog, etc.), date, page(s), volume-issue number(s), publisher, city and/or country where published	T ²
cc	E	D. WANG, ET AL.; "Toward a third generation genetic map of the human genome based on bi-allelic polymorphisms"; Abstract; Whitehead Institute/MIT Center for Genome Research, Cambridge, MA 02139 and Affymetrix, Inc., Santa Clara, CA; XP 002050641 (1996)	
	F	N.J. SCHORK, ET AL.; "Linkage disequilibrium mapping for quantitative traits within case/control settings"; Abstract; Dept. of Epidemiology and Biostatistics, Case Western Reserve University; Harvard University; The Jackson Laboratory; Genet; CEPH; XP-002089399 (1997)	
	G	EUGENE I. DRIGALENKO, ET AL.; "False Discoveries in Genome Scanning"; Genetic Epidemiology 14:779-784 (1997); 1997 Wiley-Liss, Inc.	
	H	LAURENT EXCOFFIER, ET AL.; "Maximum-Likelihood Estimation of Molecular Haplotype Frequencies in a Diploid Population"; Mol. Biol. Evol. 12(5):921-927, 1995; 1995 The University of Chicago; 0737-4038/95/1205-0021	
	I	LAURENT EXCOFFIER, ET AL.; "Incorporating Genotypes of Relatives into a Test for Linkage Disequilibrium"; Am. J. Hum. Genet. 62:171-180, 1998; The American Society of Human Genetics; 0002-9297/98/6201-0025	
	J	A. P. DEMPSTER, ET AL.; "Maximum Likelihood from Incomplete Data via the EM Algorithm"; Harvard University and Educational Testing Service; Royal Statistical Society; December 8th, 1976; pgs. 1-38	
	K	M. E. HAWLEY, ET AL.; "HAPLO: A Program Using the EM Algorithm to Estimate the Frequencies of Multi-site Haplotypes"; Dept. of Genetics, Yale University School of Medicine, The Journal of Heredity 1995:86(5)	
	L	T. SCHWEDER, ET AL.; "Plots of P-values to evaluate many tests simultaneously"; Biometrika (1982), 69, 3, pp. 493-502; Great Britain	
	M	D. C. THOMAS, ET AL.; "The Problem of Multiple Inference in Studies Designed to Generate Hypotheses"; American Journal of Epidemiology; 1985 The Johns Hopkins University School of Hygiene and Public Health; Vol 122, No. 6, pgs. 1080-1094	
	N	EDWARD I. GINNS, ET AL.; "A genome-wide search for chromosomal loci linked to mental health wellness in relatives at high risk for bipolar affective disorder among the Old Order Amish"; Proc. Natl. Acad. Sci. USA; Vol. 95, pp. 15531-15536, December 1998, Genetics	
	O	ESTER M. M. VAN LIESHOUT, ET AL.; "Polymorphic Expression of the Glutathione S-Transferase P1 Gene and Its Susceptibility to Barrett's Esophagus and Esophageal Carcinoma"; Cancer Research 59, 586-589, February 1, 1999, pgs. 586-589	

Examiner Signature		Date Considered	8/21/03
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CPK	P	GRIFFITHS, ET AL.; "An Introduction to Genetic Analysis"; 1993, U.S. XP002136574; "The Chi-Squared Test", pg. 132-135; W. H. Freeman and Company, New York	
	Q	LANDER, ET AL.; "MAPMAKER: An Interactive Computer Package of Constructing Primary Genetic Linkage Maps of Experimental and Natural Populations"; Genomics 1, 174-181 (1987); 0888-7543/87; Academic Press, Inc.	
	R	LANDER, ET AL.; "Mapping Mendelian Factors Underlying Quantitative Traits Using RFLP Linkage Maps"; Genetics 121: 185-199, January 1989; Genetics Society of America	
	S	M. XIONG, ET AL.; "Biallelic Markers in Genetics Studies of Human Diseases: Their Power, Accuracy, and Density in Population-based Linkage Analyses"; American Journal of Human Genetics, U.S., New York, NY, vol. 61, no. 4, Suppl, 1997, page 1759; XP002119235; ISSN: 0002-9297	
	T	MORELL, ET AL.; "Analysis of short tandem repeat (STR) allele frequency distributions in a Balinese population"; Hum. Mol. Gen.; vol. 4, No. 1, 1995, pgs. 85-91; XP002136573	
	U	JONATHAN CORNELIUS; "Heritabilities and additive genetic coefficients of variation in forest trees"; Canadian Journal of Forest Research, 1994, Vol. 24, pg. 372-379; XP002136575	
	V	LEONID KRUGLYAK; "The use of a genetic map of biallelic markers in linkage studies"; Nature Genetics, Vol. 17, September 1997; pgs. 21-24	
	W	JOHN G. K. WILLIAMS, ET AL.; "DNA polymorphisms amplified by arbitrary primers are useful as genetic markers"; Nucleic Acids Research, Vol. 18, No. 22; pgs. 6531-6535 (1990)	

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